	1	CLAIMS
	2	What is claimed is:
	_3	
J. W		Claim 1 . A biopolymer marker selected from the group
، عزا	5	consisting of sequence ID VDVIPVNLPGEHGQR,
	6	(R) FLATTPNSLLV WQPPR(A), HQLYIDETVNSNIPTNLR,
	7	RVDVIPVNLPGEHGQRL, SSPVVIDASTAIDAPSNLR, IHLISTQSAIPYALR or
	8	at least one analyte thereof useful in indicating at least
	9	one particular disease state.
<i>‡</i> ==4,	10	
ű	11	Claim 2. The boopolymer marker of claim 1 wherein
	11 12 13 14	said disease state is predictive of Alzheimers disease.
	_13	
H	14	Claim 3. A method for evidencing and categorizing at
the first three trees for the third the third the third three the three	15	least one disease state comprising:
	16	obtaining a sample from a patient;
	17	conducting mass spectrometric analysis on said
	18	sample;
	19	evidencing and categorizing at least one biopolymer
	20	marker sequence or analyte thereof isolated from said
	21	sample; and,
٠	22	comparing said at least one isolated biopolymer
	23	marker sequence or analyte thereof to the biopolymer

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marker sequence as set forth in claim 1;

2	marker and said biopolymer marker sequence as set forth in
3	claim 1 evidences and categorizes said at least one
4	disease state.
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6	Claim 4. The method of claim 3, wherein said step
7	of evidencing and categorizing is particularly directed to
8	biopolymer markers or analytes thereof linked to at least
9	one risk of disease development of said patient.
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11	Claim 5. The method of claim 3, wherein said step
12	of evidencing and categorizing is particularly directed to
13	biopolymer markers or analytes thereof related to the
14	existence of a particular disease state.
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16	Claim 6. The method of claim 3, wherein the sample
17	is an unfractionated body fluid or a tissue sample.
18	
19	
20	Claim 7. The method of claim 3, wherein said sample
21	is at least one of the group consisting of blood, blood
22	products, urine, saliva, cerebrospinal fluid, and lymph.
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Claim 8. The method of claim 3, wherein said mass

wherein correlation of said isolated biopolymer

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spectrometric analysis is selected from the group 1 2 consisting of Surface Enhanced Laser Desorption Ionization 3 (SELDI) mass spectrometry (MS), Maldi Qq TOF, MS/MS, TOF-TOF, and ESI-Q-TOF or an ION-TRAP. 4 5 6 Claim 9. The method of claim 3, wherein said 7 patient is a human. 8 9 A diagnostic assay kit for determining Claim 10. 10 the presence of the biopolymer marker or analyte thereof of claim 1 comprising: at least one biochemical material which is capable of specifically binding with a biomolecule which includes at least said biopolymer marker or analyte thereof, and means for determining binding between said biochemical material and said biomolecule; whereby at least one analysis to determine a presence 18 of a marker, analyte thereof, or a biochemical material 19 specific thereto, is carried out on a sample. 20 21 The diagnostic assay kit of claim 10, 22 wherein said biochemical material or biomolecule is 23 immobilized on a solid support.

	i	Claim 12. The diagnostic assay kit of claim 10
	2	including:
	3	at least one labeled biochemical material.
	4	
	5	Claim 13. The diagnostic assay kit of claim 10,
	6	wherein said biochemical material is an antibody.
	7	
	8	Claim 14. The diagnostic assay kit of claim 12,
	9	wherein said labeled biochemical material is an antibody.
	10	
	11	Claim 15. The diagnostic assay kit of claim 10,
	12	wherein the sample is an unfractionated body fluid or a
_	13	tissue sample.
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ă.	15	Claim 16. The diagnostic assay kit of claim 10,
	16	wherein said sample is at least one of the group
1	17	consisting of blood, blood products, urine, saliva,
	18	cerebrospinal fluid, and lymph.
	19	
	20	Claim 17. The diagnostic assay kit of claim 10,
	21	wherein said biochemical material is at least one
	22	monoclonal antibody specific therefore.
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Claim 18. \setminus A kit for diagnosing, determining risk-

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assessment, and identifying therapeutic avenues related to
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   2
        a disease state comprising:
             at least\ one biochemical material which is capable of
   4
        specifically binding with a biomolecule which includes at
        least one biopolymer marker selected from the group
   5
   6
        consisting of sequence ID VDVIPVNLPGEHGQR,
   7
        (R) FLATTPNSLLVSWQPPR(A), HQLYIDETVNSNIPTNLR,
        RVDVIPVNLPGEHGQRL, $SPVVIDASTAIDAPSNLR, IHLISTQSAIPYALR or
   8
        analyte thereof related to said disease state; and
             means for determining binding between said
        biochemical material and said biomolecule;
  11
  12
             whereby at least one analysis to determine a presence
        of a marker, analyte thereof, or a biochemical material
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5 14
        specific thereto, is carried out on a sample.
<del>1</del> 15
TU 16
                        The kit of claim 18, wherein said
             Claim 19.
17
        biochemical material or biomolecule is immobilized on a
  18
        solid support.
  19
  20
             Claim 20. The kit of claim 18 including:
  21
             at least one labeled biochemical material.
  22
  23
                          The kit of claim 18, wherein said
             Claim 21.
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biochemical material is an antibody.

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	4	Claim 23. The kit of claim 18, wherein the sample is
	5	an unfractionated body fluid or a tissue sample.
	6	
	7	Claim 24. The kit of claim 18, wherein said sample
	8	is at least one of the group consisting of blood, blood
	9	products, urine, saliva, cerebrospinal fluid, and lymph.
energ.	10	
u u	11	Claim 25. The kit of claim 18, wherein said
HELL ENDE	.12	biochemical material is at least one monoclonal antibody
	13	specific therefore.
	14	
	15	Claim 26. The kit of claim 18, wherein said
The straightfun from the fire	16	diagnosing, determining risk assessment, and identifying
	17	therapeutic avenues is carried out on a single sample.
	18	
	19	Claim 27. The kit of claim 18, wherein said
	20	diagnosing, determining risk assessment, and identifying
	21	therapeutic avenues is carried out on multiple samples
	22	such that at least one analysis is carried out on a first

sample and at least another analysis is carried out on a

The kit of claim 20, wherein said labeled

second sample.

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Claim 22.

biochemical material is an antibody.

The kit of claim 27, wherein said first 1 Claim 28. and second samples are obtained at different time periods. 3 Polyclonal antibodies produced against a Claim 29. marker sequence ID selected from the group consisting of sequence VDVIPVNLPGEHGQR, (R) FLATTPNSLLVSWQPPR(A), HQLYIDETVNSNIPTNLR, RVDVIPVNLPGEHGQRL, 7 SSPVVIDASTAIDAPSNLR, IHLISTQSAIPYALR or at least one 8 9 analyte thereof in at least one animal host. 10 0 0 11 Claim 30. An antibody that specifically binds a □ 12 biopolymer including \a marker selected from the group consisting of sequence ID VDVIPVNLPGEHGQR, 13 **T** 14 (R) FLATTPNSLLVSWQPPR (A) \ HQLYIDETVNSNIPTNLR, RVDVIPVNLPGEHGQRL, SSPVVIDASTAIDAPSNLR, IHLISTQSAIPYALR 15 or at least one analyte thereof. Claim 31. The antibody of claim 30 that is a 18 19 monoclonal antibody. 20 21 Claim 32. The antibody of claim 30 that is a 22 polyclonal antibody. 23 A process for identifying therapeutic Claim 33.

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2 conducting an analysis as provided by the kit of claim 18; and 3 4 interacting with a biopolymer selected from the group consisting of sequence ID VDVIPVNLPGEHGQR, 5 (R) FLATTPNSLLVSWQPPR(A), HQLYIDETVNSNIPTNLR, 6 RVDVIPVNLPGEHGQRL, SPVVIDASTAIDAPSNLR, IHLISTQSAIPYALR 7 or at least one analyte thereof; 8 whereby therapeutic avenues are developed. The process for identifying therapeutic Claim 34. 11 avenues related to a disease state in accordance with 12 claim 33, wherein said therapeutic avenues regulate the 13 **T** 14 presence or absence of the biopolymer selected from the 15 group consisting of sequence ID VDVIPVNLPGEHGQR, N 16 (R) FLATTPNSLLVSWQPPR(A), HQ14YIDETVNSNIPTNLR, **17** RVDVIPVNLPGEHGQRL, SSPVVIDASTAIDAPSNLR, IHLISTQSAIPYALR or at least one analyte thereof. 18 20 Claim 35. The process for identifying therapeutic 21 avenues related to a disease state in accordance with

avenues related to a disease state comprising:

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- 23 include at least one avenue selected from a group
- 24 consisting of 1)utilization and recognition of said

claim 33, wherein said therapeutic avenues developed

1 biopolymer markers, variants or moieties thereof as direct 2 therapeutic modalities, either alone or in conjunction 3 with an effective amount of a pharmaceutically effective carrier; 2) validation of therapeutic modalities or disease 4 5 preventative agents as a function of biopolymer marker 6 presence or concentration; 3) treatment or prevention of a 7 disease state by formation of disease intervention 8 modalities; 4) use of biopolymer markers or moieties 9 thereof as a means of elucidating therapeutically viable 10 agents, 5) instigation of a therapeutic immunological response; and 6) synthesis of molecular structures related to said biopolymer markers, moieties or variants thereof 13 which are constructed and arranged to therapeutically intervene in said disease state. Claim 36. The process for identifying therapeutic

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avenues related to a disease state in accordance with claim 35, wherein said treatment or prevention of a disease state by formation of disease intervention modalities is the formation of biopolymer/ligand conjugates which intervene at receptor sites to prevent, delay or reverse a disease process.

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24 The process for identifying therapeutic Claim 37.

- ·
- 1 avenues related to a disease state in accordance with
- 2 claim 35, wherein said means of elucidating
- 3 therapeutically viable agents includes use of a
- 4 bacteriophage peptide display library or a bacteriophage
- 5 antibody library.

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Claim 38. A process for regulating a disease state by controlling the presence or absence of a biopolymer selected from the group consisting of sequence ID

- 10 VDVIPVNLPGEHGQR, (R) FLATTPNSLLVSWQPPR(A),
- 11 HQLYIDETVNSNİPTNLR RVDVIPVNLPGEHGQRL,
- 12 SSPVVIDASTAIDAPSNLR, IHLISTQSAIPYALR or at least one
- 13 analyte thereof.

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